U.S. DEPARTMENT OF COMMERCE ATTORNEY DOCKET NO. **FORM PTO-1390** 100564-00107 PATENT AND TRADEMARK OFFICE (REV 5-93) DATE: March 11, 2002 TRANSMITTAL LETTER TO THE UNITED STATES DESIGNATED/ELECTED OFFICE (DO/EO/US) U.S. APPLN, NO. CONCERNING A FILING UNDER 35 U.S.C. 371 (IF KNOWN, SEE 37 C F.R. New 10/0699 INTERNATIONAL FILING DATE PRIORITY DATE CLAIMED INTERNATIONAL APPLICATION NO. September 8, 2000 September 10, 1999 PCT/EP00/08812 TITLE OF INVENTION: ANT-1 AS DRUG TARGET APPLICANT(S) FOR DO/EO/US: Stefan GRIMM, Manuel BAUER, Alexis SCHUBERT and Ursula CRAMER 1. This is a FIRST submission of items concerning a filing under 35 U.S.C. 371. (THE BASIC FILING FEE IS ATTACHED) 2. This is a SECOND or SUBSEQUENT submission of items concerning a filing under 35 U.S.C. 371. This express request to begin national examination procedures [35 U.S.C. 371(f)] at any time rather than delay examination until the expiration of the applicable time limit set in 35 U.S.C. 371(b) and PCT Articles 22 and 39(1). A proper demand for International Preliminary Amendment was made by the 19th month from the earliest claimed priority date. A copy of the International Application as filed [35 U.S.C. 371(c)(2)] a.
is transmitted herewith (required only if not transmitted by the International Bureau). has been transmitted by the International Bureau. b. 🖾 is not required, as the application was filed in the United States Receiving Office (RO/US). 6. A translation of the International Application into English [35 U.S.C. 371(c)(2)]. are transmitted herewith (required only if not transmitted by the International Bureau). a. 🗍 have been transmitted by the International Bureau. have not been made; however, the time limit for making such amendments has NOT expired. d. A have not been made and will not be made. 8.
A translation of the amendments to the claims under PCT Article 19 [35 U.S.C. 371(c)(3)]. 9. An oath or declaration of the inventor(s) [35 U.S.C. 371(c)(4)]. 10. A translation of the annexes to the International Preliminary Examination Report under PCT Article 36 [35 U.S.C. 371(c)(5)]. Items 11 - 16 below concern other document(s) or information included: 11. An Information Disclosure Statement under 37 C.F.R. 1.97 and 1.98. A FIRST preliminary amendment. 13. A SECOND or SUBSEQUENT preliminary amendment. 14. A substitute specification.

15. A change of power of attorney and/or address letter.

Drawings (⊠ FIGS 1-8. 8 sheets)

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Other items or information: PCT/ISA/210 and PCT/IPEA/416.

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PATENT APPLICATION

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re the Application of:

GRIMM et al.

Group Art Unit: Unknown

Application No.: New U.S. Patent Application

Examiner: Unknown

Filed: March 11, 2002

Attorney Dkt. No.: 100564-00107

For: ANT-1 AS DRUG TARGET

PRELIMINARY AMENDMENT

Commissioner for Patents Washington, D.C. 20231

March 11, 2002

Sir:

Prior to initial examination of the application, please amend the above-identified application as follows:

IN THE CLAIMS:

Please cancel claims 1-31 without Prejudice or disclaimer.

- 32. A method for the inhibition of apoptosis, comprising contacting a cell with an effective amount of a substance capable of inhibiting the activity of adenine nucleotide translocase-1 (ANT-1).
- 33. The method of claim 32, wherein said cell is a mammalian cell.
- 34. The method of claim 33, wherein said cell is associated with a pathogenic disorder.

- 35. The method of claim 32, wherein the activity of ANT-1 is inhibited on the nucleic acid level.
- 36. The method of claim 35, wherein the inhibition is effected by reducing ANT-1 gene expression.
- 37. The method of claim 36, wherein the activity of the endogenous ANT-1 promoter is reduced.
- 38. The method of claim 32, wherein the activity of ANT-1 is inhibited on the protein level.
- 39. The method of claim 38, wherein the inhibition is effected by adding ANT-1 protein antagonists.
- 40. The method of claim 39, wherein the antagonist is cyclophilin D.
- 41. The method of claim 32, wherein an apoptosis-inducing signal transduction pathway is inhibited, said pathway being activated by ANT-1.
- 42. A method for the treatment of diseases associated with excessive apoptosis, comprising the step of administering to a subject in need thereof a pharmaceutically effective amount of a substance capable of inhibiting the activity of adenine nucleotide translocase (ANT-1).
- 43. The method of claim 42, wherein the disease is a degenerative disease.
- 44. The method of claim 43, wherein the disease is dilated cardiomyopathy.

- 45. A method for identifying substances suitable for apoptosis inhibition comprising the step of determining the capability of a test substance to inhibit the activity of ANT-1.
- 46. The method of claim 45, wherein the capability of a test substance to bind ANT-1 or a domain thereof is determined.
- 47. The method of claim 45, wherein the capability of a test substance to bind the N-terminal domain of ANT-1 is determined.
- 48. The method of claim 45, wherein the capability of a test substance to inhibit the binding of ANT-1 to natural binding partners thereof is determined.
- 49. The method of claim 45, which is carried out as a high-throughput assay.
- 50. The method of claim 49, comprising a parallel determination of at least 96 test compounds.
- 51. The method of claim 45, which is carried out as a cell-based assay.
- 52. The method of claim 50, which is carried out as an assay using ANT-1-containing cell fractions or ANT-1-containing whole cells.
- 53. The method of claim 45, which is carried out as a molecular-based assay using an isolated protein selected from ANT-1 or a domain thereof.
- 54. The method of claim 43, wherein a recombinant protein is used.
- 55. The method of claim 45, wherein the determining step comprises the measurement of apoptosis induction.

REMARKS

Claims 32-62 are pending in this application. By this Amendment, claims 1-31 have been cancelled. No new matter is contained in the amendments.

Please charge any fee deficiency or credit any overpayment to Deposit Account No. 01-2300.

Respectfully submitted,

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ANT-1 as Drug Target

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Description

The present invention refers to the use of adenine nucleotide translocase-1 (ANT-1), a central component of the permeability transition pore in mitochondria, as a drug target, particularly for the treatment of dilated cardiomyopathy. Further, an assay for the detection of pharmacologically active substances is disclosed which inhibit ANT-1 activity. In addition, ANT-1 expression is claimed as a diagnostic marker for dilated cardiomyopathy.

Apoptosis is a form of cell death that plays a role in development, tissue homeostasis and disease (White, 1996). Its induction must be tightly regulated. Otherwise, serious consequences may follow: A hyperactive apoptosis induction might lead to degenerative diseases like Alzheimer's disease (Loo et al., 1993); a reduced activity can contribute to the multistep process of tumorigenesis, since tumor cells are exposed to multiple proapoptotic stimuli (McGill, 1997). The induction of apoptosis is therefore governed by an elaborate array of checks and balances in the cell. Eventually, a family of cysteine proteases, the so-called caspases, is activated (Salvesen and Dixit, 1997). These enzymes can cleave specific substrates in the cell which leads to the typical apoptotic phenotype and the self-destruction of the cell.

The first indication that mitochondria play a role in apoptosis induction was the observation that an in vitro system for apoptosis induction required the presence of mitochondria (Newmeyer et al., 1994). The permeability transition pore (PT) complex (Zoratti and Szabo, 1995), a protein aggregate that resides in contact sites of the inner and outer mitochondrial membrane, was subsequently identified as being responsible for apoptosis induction

(Petit et al., 1995; Zamzami et al., 1995b). Multiple pharmacological stimuli have been shown to activate this complex by as yet unknown means (Fulda et al., 1998; Petit et al., 1995; Zamzami et al., 1998). The activated complex leads to the collapse of the potential over the inner mitochondrial membrane ($\Delta\Psi_m$), swelling of mitochondria and the generation of oxygen radicals (Vander Heiden et al., 1997; Zamzami et al., 1995a). The subsequent release of apoptogenic caspases (Susin et al., 1999a) and a putative oxidoreductase (Susin et al., 1999b) aids in apoptosis induction.

Many genes involved in apoptosis have the dominant capacity to induce cell death upon overexpression. This feature is even conserved across species (McCarthy and Dixit, 1998). It might be explained by the fact that the overexpressed proteins engage in protein-protein contacts and can thereby create the signal for apoptosis (Yang et al., 1998). Consequently, we have recently developed a screen for dominant, apoptosis-inducing genes (Grimm and Leder, 1997). The screen is based on the iterative transfection of small plasmid pools of a normalized cDNA library into mammalian cells and the morphological determination of apoptosis induction.

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Here we describe the isolation of adenine nucleotide translocase-1 (ANT-1), a central component of the permeability transition pore, using such a screen. ANT-1 has recently been shown to be required for apoptosis mediated by Bax, another component of the PT complex (Marzo et al., 1998a). Conversely, ANT-1 can induce apoptosis in a Bax-dependent manner. For this, ANT-1 must be activated pharmacologically by the chemical attractyloside which arrests ANT-1 in a specific conformation and causes PT pore opening (Marzo et al., 1998a). In our experiments with ANT-1, however, such a secondary signal is not required for apoptosis induction. ANT-1's apoptosis activity does not seem to depend on its known function for ADP/ATP exchange because several transport inactive mutants could still lead to cell death. Surprisingly, we found that a very homologous protein, the ANT-2 transporter, was inactive for apoptosis

induction. Furthermore, ANT-1 could not elicit a form of cell death in yeast. This is in contrast to Bax that directly interacts with ANT-1 and can dominantly induce cell death in yeast. This result suggests that ANT-1mediated apoptosis induction depends on protein-protein interactions that are specific for mammalian cells and is independent of Bax-mediated apoptosis. Since ANT-1 is activated by overexpression for apoptosis induction, it is interesting to note that this gene is already highly expressed in mitochondria. This suggests that in a normal cell it must be kept inactive by other proteins of the PT pore which implies important stoichiometric correlations between the various components of this complex. Consistent with this, we found that cyclophilin D, another component of the PT pore can repress ANT-1-induced apoptosis. Furthermore, our data help to explain the observed apoptosis induction in dilated cardiomyopathy (DCM), a degenerative disease of the heart muscle which is marked by a dramatic increase of the expression level of ANT-1 and by excessive apoptosis induction.

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These findings are up to now the most direct genetic evidence that the PT pore can signal apoptosis and may lead to the molecular elucidation of how this complex can be activated for apoptosis induction.

A first aspect of the present invention is a method for the inhibition of apoptosis comprising the step of contacting a cell which is susceptible to apoptosis with an effective amount of a substance capable of inhibiting the activity of adenine nucleotide translocase-1 (ANT-1), particularly the apoptosis-inducing activity of ANT-1.

The cell is preferably a vertebrate cell, more preferably a mammalian cell and, still more preferably, a human cell, e.g. a human myocyte.

The inhibition may effect an apoptosis-inducing signal transduction pathway, said pathway being activated by ANT-1. The activity of ANT-1

may be inhibited on the nucleic acid level and/or on the protein level. Inhibition on the nucleic acid level may be effected by reducing ANT-1 gene expression, e.g. by using antisense-oligonucleotides which may inhibit the ANT-1 gene transcription, by using ribozymes capable of cleaving ANT-1 transcripts or by inhibiting the activity of the endogenous promoter, e.g. via ANT-1-specific transcription factors. The reduction of ANT-1 expression may also be achieved by influencing the activity of proteins that act upstream of these transcription factors, e.g. such as protein kinases and phosphatases. Inhibition on the protein level may be effected by anti-ANT-1 antibodies, by ANT-1 regulatory proteins, e.g. by proteins occurring in permeability transition pores such as cyclophilin D or by low molecular weight substances which are capable of inhibiting the apoptotic or proapoptotic activity of ANT-1.

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A further subject matter of the present invention is a method for the treatment of diseases associated with excessive apoptosis, comprising the step of administering to a subject in need thereof a pharmaceutically effective amount of a substance capable of inhibiting the activity of adenine nucleotide translocase-1 (ANT-1), particularly the apoptosis-inducing activity of ANT-1. Examples of diseases which may be treated by administering ANT-1 antagonists are dilated cardiomyopathy and other human pathogenic disorders associated with excessive apoptosis such as degenerative diseases. More preferably, the disease is dilated cardiomyopathy.

The ANT-1 antagonist may be administered as a pharmaceutical composition comprising the active agent, optionally together with pharmaceutically acceptable diluents, carriers or adjuvants. The pharmaceutical composition may be suitable for oral, parenteral, e.g. intradermal, intravenous or intramuscular, rectal, nasal and topical applications. The composition may be an injectable solution, ointment, cream or spray. Further, the composition may have retardation properties, e.g. showing a delayed release of the active agent.

The dosage of the active agent depends on the specific compound being administered, the type and the severity of the disease. For example, a dosage from 0.01 mg to 100 mg per day and per kg body weight of the active agent is suitable.

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Still a further aspect of the present invention is a method for identifying substances suitable for apoptosis inhibition, comprising the step of determining the capability of a test substance to inhibit the activity of ANT-1, particularly the apoptosis-inducing activity of ANT-1. Preferably, the capability of a test substance to bind ANT-1 or a domain thereof may be determined. Particularly, the N-terminal domain of ANT-1 comprising amino acids 1-150, preferably amino acids 1-200, is suitable for determining the binding of test substances. Alternatively, the capability of test substances to inhibit the binding of ANT-1 to natural binding partners thereof such as cyclophilin D may be determined.

throughput assay which preferably comprises a parallel determination of at least 24 and, more preferably, at least 96 test compounds. The assay may be carried out as a cell-based assay, wherein a test system is used comprising ANT-1-containing cell fractions or ANT-1-containing whole cells. Alternatively, the assay may be carried out as a molecular-based assay comprising a substantially purified and isolated protein selected from ANT-1 or a domain thereof, particularly an N-terminal domain thereof. The substantially purified ANT-1 protein may be a recombinant ANT-1 protein which may a soluble protein (by deletion of membrane anchor domains) or

The identification of ANT-1 antagonists may be carried out as a high-

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a membrane-bound protein.

In a cell-based assay the determining step may comprise the measurement of apoptosis induction. Apoptosis induction may be measured by a parameter which is associated with apoptosis such as DNA fragmentation, caspase activation or characteristic alterations in cell morphology or other

qualitative and/or quantitative methods of apoptosis measurement known in the field. In molecular-based assay systems the determination may comprise the measurement of protein-protein interactions, e.g. the interaction of ANT-1 with other proteins, or direct measurement of the interaction of test compounds with ANT-1.

As the invention describes upregulated ANT-1 as a drug target in dilated cardiomyopathy or other degenerative diseases, a further aspect of this invention is the examination of the ANT-1 expression level in patients as a diagnostic marker. The ANT-1 expression can be detected and quantified by conventional methods, e.g. on the RNA level by hybridization and/or amplification procedures or on the protein level by antibody and/or activity assays. Thus, the invention relates to a method for the diagnosis of a degenerative disease or a predispositon therefor comprising detecting the ANT-1 expression in a sample from tissue and/or body fluids of a subject to be tested, wherein elevated ANT-1 expression is indicative for a degenerative disease or a predisposition therefor.

The present invention is further illustrated by the following Figures and Examples.

Description of Figures

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- Fig. 1. Adenine nucleotide translocase-1 (ANT-1) expression leads to phenotypic apoptosis induction. The empty vector or an expression plasmid for ANT-1 were transiently transfected into 293T cells. After 16 hours, phase contrast pictures were taken at a 200-fold magnification.
- Fig. 2. Effect of ANT-1 expression on mitochondria. (A) ANT-1 overexpression leads to the collapse of the inner mitochondrial membrane potential. HeLa cells were transfected with an expression vector for GFP (2 μ g) together with a control vector ("Vector"; 2 μ g) or a plasmid for

ANT-1 ("ANT-1"; $2~\mu g$). After 16 hours, the cells were treated with CMXRos which stains mitochondria with an intact membrane potential. Subsequently, phase contrast and fluorescence microscopy pictures for GFP and CMXRos activity were taken. (B) ANT-1 induces the release of cytochrome c from mitochondria. 293T cells were transfected with a control vector or an expression vector for ANT-1. Cytoplasmic extracts were prepared and assayed for the presence of cytochrome c by a western blot. Molecular weight standards are given on the left. Equal loading of the gel is indicated by an upper unspecific band with equal intensity.

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Fig. 3. Effect of ANT-1 on protein and DNA degradation. (A) ANT-1 leads to the degradation of poly-ADP-ribose polymerase (PARP). 293T cells were transfected with an empty control vector, an expression vector for the "death domain" protein RIP or with ANT-1. 16 hours later, nuclear extracts of the transfected cells were prepared and investigated for the status of PARP in a western blot. The generated PARP fragment is indicated by an arrow. An unspecific signal (o) served as an internal control for equal loading of the gel. (B) Inhibition of ANT-1 apoptosis by a specific inhibitor for caspases. ANT- 1 (1 μ g) and a control vector (10 μ g) or an expression vector (10 μ g) for the caspase inhibitor p35 from baculovirus were cotransfected, and the specific apoptosis induction was determined by FACS analysis. The means and the standard deviations are indicated (n=3) (C) ANT-1 induces internucleosomal DNA cleavage. 293T cells were transfected with an empty vector or with an expression construct for ANT-1. Low molecular weight DNA was isolated from transfected cells, separated on a 2% agarose gel and stained with ethidium bromide.

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Fig. 4. Mutational analysis of ANT-1's apoptosis activity (A) Cell death induction by point mutants of ANT-1. Wild type ANT-1 (ANT-1 WT) and six point mutants that have been shown to be deficient for ADP/ATP transport were transfected into 293T cells. The WT amino acid, its position as well as the mutated residue are given for each mutant construct. After 16

hours, apoptosis induction by the various constructs was measured by an ELISA for internucleosomal DNA fragments. The DNA fragmentation as percentage of control is given as an index for apoptosis induction. The means and the standard deviations are indicated (n=3) (B) Apoptosis activity of ANT-1 deletion mutants. Wild type ANT-1 (1 μ g) and three C-terminal deletion mutants (1 μ g) were transiently transfected into 293T cells. The last C-terminal residue of each deletion mutant is indicated. After 16 hours, the constructs were tested for apoptosis induction by a specific ELISA as described in A.

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Fig. 5. Effect of ANT-1 and ANT-2 on transfected cells. (A) Comparison between the apoptosis capacities of ANT-1 and ANT-2. Expression plasmids for ANT-1 (1 μ g) and ANT-2 (1 μ g) were transfected in 293T cells. After 16 hours apoptosis induction in transfected cells was measured by FACS analysis of sub-G1 positive cells. The specific apoptosis induction above background is shown as percentage of apoptotic cells relative to all transfected cells. The means and the standard deviations are indicated (n=3). (B) Expression and localization of ANT-1 and ANT-2. 293T cells were transfected with a control vector or expression vectors for HA-tagged ANT-1 and ANT-2. Mitochondrial extracts of transfected cells were prepared and investigated for the presence of the proteins with an anti-HA antibody in a western blot. Equal loading of the gel was verified by two unspecific upper bands.

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Fig. 6. Effect of Bax and ANT-1 expression on the growth of yeast cells. Bax and ANT-1 cloned in expression vectors under the control of a galactose and raffinose-inducible promoter and the empty control vector were introduced into yeast cells. 4,5x10⁴ yeast cells of three independent clones each were plated on glucose (glu)- or galactose and raffinose (gal/raf)-containing agar. The growth of the yeast cells was monitored 36 hours later.

Fig. 7. Inhibition of ANT-1 induced apoptosis. (A) ANT-1-induced apoptosis can be repressed by bongkrekic acid, a specific inhibitor of the PT pore. An expression vector for ANT-1 (1 μ g) was transfected into plates with 293T cells. One group of plates was left untreated, the other was supplied with 50 μ M bongkrekic acid ("BA") for 16 hours after the transfection. Apoptosis was quantified as in figure 5. (B) ANT-1 apoptosis can be repressed by cyclophilin D, another component of the permeability transition pore. ANT-1 (1 μ g) and an expression plasmid (10 μ g) for cyclophilin D ("Cyclo D") or a control vector (10 μ g) were transfected into 293T cells, and apoptosis was determined as described in figure 5.

Examples

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1.Materials and methods

1.1 Plasmids

The point mutants of ANT-1 were engineered by recombinant PCR using suitable primers. For the PCR reaction, Pwo, a thermostable polymerase with proof-reading function (Roche Diagnostics, Mannheim, Germany) was employed. All amino acid changes were verified by sequencing. Deletion mutagenesis of ANT-1 was achieved with PCR using the same enzyme. ANT-2 was amplified from a mouse kidney cDNA library, cyclophilin D from a 293T library, both by using specific primers. The correct sequence was verified by sequencing. The eukaryotic expression vector for baculovirus p35 has been described (Clem and Miller, 1994).

1.2 Immuno Blotting

For detecting cytochrome c, 293T cells were transfected with an expression vector for ANT-1. After 18 hours the cells were harvested and cytoplasmic extracts isolated as described (Ferrari et al., 1998). Protein samples of 30

 μ g were loaded on SDS-polyacrylamide gels (10%) and then electrophoretically transferred onto PVDF-membranes (Amersham, Braunschweig, Germany). Western blots were probed with a monoclonal cytochrome c antibody (Pharmingen, San Diego, CA) and anti-mouse-horseradish-peroxidase conjugated antibody prior to ECL-based detection (Amersham). For the poly-ADP-ribose polymerase (PARP) western blot, nuclear fractions were obtained by differential centrifugation as described previously (Schreiber et al., 1989). Aliquots of 50 μ g protein were subjected to SDS-PAGE (8% polyacrylamide) and blotted onto PVDF-membranes. PARP and cleavage products were detected using a polyclonal serum directed against full length PARP (Roche Diagnostics), anti-rabbit-horseradish-peroxidase conjugated antibody (Amersham, Braunschweig) and the ECL-system. For the detection of HA-tagged ANT-isoforms, mitochondria were isolated by differential centrifugation as described previously (Brustovetsky and Klingenberg, 1996). 30 μ g protein samples were separated on 10% polyacrylamide gels and blotted onto PVDF-membranes. Detection was performed using a mouse monoclonal antibody raised against the influenza virus HA-peptide (Roche Diagnostics, Mannheim, Germany) and the ECL-system (Amersham, Braunschweig).

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1.3 Yeast methods

Standard yeast methods were applied (Ausubel et al., 1991). For the inducible expression of Bax and ANT-1, the cDNAs were subcloned into pYESTrp2 (Invitrogen) in which the B42-fusion moiety was removed.

1.4 Apoptosis detection

Low molecular weight DNA from apoptotic cells was isolated and detected as described (Grimm and Leder, 1997). Apoptosis induction was measured by an ELISA (Roche Diagnostics) that is specific for nucleosomal DNA fragments that are released during apoptosis. The recommendations of the

manufacturer were followed. Equal transfection efficiencies were monitored by cotransfecting an expression plasmid for GFP. Other apoptosis quantifications were performed by flow cytometry (Bitzer et al., 1999). A cotransfected GFP expression plasmid was used to assess the transfection efficiency. The apoptotic population was put in relation to the percentage of GFP-positive cells. The apoptotic background of the vector-control was subtracted to obtain the specific apoptosis induction. Each condition was tested in at least three independent experiments.

1.5 Mitochondrial membrane gradient detection

After transfecting an expression vector for ANT-1 together with a plasmid for GFP, cells were loaded with Mitotracker RedCMXRos (Molecular probes, OR, USA) according to the suggestions of the manufacturer. Images were documented using a Zeiss Axioplan fluorescence microscope (Zeiss Oberkochen, Germany). RedCMXRos fluorescence was excited at 546 nm and emission was imaged at >590 nm. GFP fluorescence was excited at 450-490 nm and emission was monitored at 515-565 nm.

2.Results

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2.1 ANT-1 induces cell death

Using our screen for dominant, apoptosis-inducing genes, we isolated a cDNA whose expression elicited an especially fast and strong apoptosis response in cells. Sequencing revealed that the gene encoded ANT-1, the ADP/ATP translocator protein of the PT pore (Marzo et al., 1998b). ANT was active in a wide variety of cell types: every cell line tested so far underwent apoptosis after stimulation by ANT-1 (not shown). Figure 1 shows the phenotype of control- and ANT-1-transfected HEK 293T cells: whereas the vector-transfected cells displayed a normal morphology, ANT-1

expressing cells showed a constricted cytoplasm and blebbing of the plasma membrane.

Since ANT-1 is a component of the PT pore whose activation leads to a dissipation of the proton gradient $\Delta\Psi_m$ over the inner mitochondrial membrane (Marzo et al., 1998b), we tested ANT-1 transfected cells for this membrane gradient. To this end, we cotransfected ANT-1 and an expression vector for the green fluorescent protein (GFP) into HeLa cells. The membrane gradient in transfected and therefore GFP-positive cells was assayed with CMXRos, a dye whose fluorescence is dependent on an intact membrane potential. Figure 2A shows that ANT-1 expressing cells displayed a reduced mitochondrial fluorescence with this dye. This differential staining is a very early feature of ANT-1-mediated apoptosis induction since it could be observed in cells which did not yet display an altered phenotype.

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One of the consequences of mitochondrial damage is the release of cytochrome c from the inner mitochondrial membrane space (Kluck et al., 1997; Yang et al., 1997). Therefore, we assayed cytoplasm of ANT-1 transfected cells for the presence of cytochrome c. Using a western blot, we could detect this protein in the cytoplasmic fraction of ANT-1 transfected cells (figure 2B). The released cytochrome c can bind Apaf-1, a mammalian homolog of the C. elegans gene ced-4 which leads to the aggregation and activation of downstream caspases (Srinivasula et al., 1998). We therefore investigated the status of poly-ADP-ribose polymerase (PARP), one of the best-known caspase substrates, in ANT-1 transfected cells. A western blot for PARP (figure 3A) revealed that ANT-1 transfected cells degraded PARP and generated the expected protein fragments. Since we have also observed apoptosis induction in MCF-7 cells (data not shown), a cell line that does not express caspase-3 (Janicke et al., 1998), we conclude that ANT-1-induced apoptosis is not dependent on this particular isoenzyme.

Bax, a member of the Bcl-2 gene family, has recently been shown to interact directly with ANT-1 (Marzo et al., 1998a) and is known to induce apoptosis (Oltvai et al., 1993). However, caspase activation does not seem to be a necessary event for Bax-induced cell death (Xiang et al., 1996). Therefore, we wanted to know whether caspase activation is required for ANT-1 apoptosis induction. To address this, we cotransfected ANT-1 and an expression vector for p35, a broad-range caspase inhibitor from baculovirus (Clem and Miller, 1994). A quantitative apoptosis assay revealed a 6-fold reduction in apoptosis induction when p35 was cotransfected (figure 3B). Another substrate for caspases is ICAD, an inhibitor of the DNase CAD (Sakahira et al., 1998). Its degradation leads to the activation of CAD and the internucleosomal degradation of the DNA, a well-known biochemical sign of apoptosis. To investigate the status of the DNA, we transfected 293T cells with a control vector or ANT-1 and isolated the low molecular weight DNA. Only ANT-1 transfected cells displayed the typical DNA ladder (figure 3C). These results show that ANT-1 leads to all phenotypic and biochemical alterations associated with apoptosis.

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2.2 The N-terminal half of ANT-1 is sufficient for apoptosis induction

ANT-1 was originally described as an ADP/ATP transporter (Riccio et al., 1975). We therefore theorized that overexpressed ANT-1 leads to an increased ADP/ATP exchange which results in a net-import of positive charges over the inner mitochondrial membrane, the subsequent collapse of the membrane potential and apoptosis induction. To assess whether the transporter activity of ANT-1 is required for its apoptosis induction, we generated six different point mutations all of which have been shown to impede ANT-1's activity to transport ADP and ATP (Muller et al., 1996; Muller et al., 1997). However, upon transfection into 293T cells we have not observed any difference relative to the wildtype (WT) in their potential to induce apoptosis as measured by the release of internucleosomal DNA

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fragments (figure 4A). ANT-1 is a member of the mitochondrial carrier family which consists of proteins with three 100 residue repeats in which alternating regions of hydrophilic and hydrophobic residues exist. As integral membrane proteins they span the membrane six times, interrupted by short amphipathic helices that are likely to form the transport active moiety (Klingenberg and Nelson, 1994). To map the domains responsible for apoptosis induction, we have created various deletion mutants of ANT-1:ANT-1 Δ 217 lacks the last transmembrane domain and the last amphipathic helix, ANT-1 Δ 142 cuts the molecule in half and comprises only the first two transmembrane domains and the first amphipathic helix. ANT-1 Δ 102 contains only the first membrane spanning domain. Figure 4B demonstrates that ANT-1 Δ 217 is as efficient for apoptosis induction as WT ANT-1, whereas ANT-1 Δ 142 revealed a 50% reduction of its apoptosis potential. The largest deletion, ANT-1 Δ 102, was inactive for apoptosis induction.

2.3 ANT-2 is unable to induce apoptosis

The PT pore complex is supposed to function by the conversion of the specific ADP/ATP exchange transporter into an unselective pore (Zoratti and Szabo, 1995). However, since over half of the ANT-1 protein is dispensable for apoptosis induction, we wanted to know whether ANT-1 induces apoptosis by forming an unspecific pore on its own. We therefore assessed the specificity of apoptosis induction by ANT-1. To this end we made use of ANT-2, an over 90% identical gene to ANT-1 that is likewise ubiquitously expressed (Doerner et al., 1997). Transfection of ANT-2 did not lead to apoptosis induction as measured by FACS analysis (figure 5A). The correct mitochondrial localization and efficient expression of ANT-2 was verified by western blot (figure 5B).

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2.4 Growth of yeast cells is not affected by ANT-1

Bax, a member of the Bcl-2 gene family, can induce a form of cell death by overexpression in yeast cells (Greenhalf et al., 1996; Jurgensmeier et al., 1997; Zha et al., 1996), probably by activating ANT-1 and the PT pore (Greenhalf et al., 1996; Marzo et al., 1998a). We therefore wanted to determine whether ANT-1 has the same capacity. To address this, we subcloned ANT-1 and Bax into a galactose and raffinose inducible yeast vector. The induction of these genes on suitable agar plates revealed that Bax led to a reduction in the growth of these cells, indicating cell death. In contrast, ANT-1 expressing yeast cells were completely unaffected and grew as well as on the control plate (figure 6).

2.5 Cyclophilin D can repress apoptosis induced by ANT-1

Our experiments with ANT-2 and cell death in yeast suggested that ANT-1's apoptotic activity is mediated by specific protein-protein interactions rather than by forming an active PT pore on its own. To check whether ANT-1 expression nevertheless activates the endogenous PT pore, we made use of bongkrekic acid, a specific inhibitor for the PT pore (Klingenberg et al., 1970). 50 μ M of this compound repressed ANT-1 induced apoptosis in a quantitative apoptosis assay more than 2-fold (figure 7A). We therefore speculated that ANT-1 might titrate out some components of the endogenous PT pore and that replenishing these proteins might repress apoptosis. Consequently, we cotransfected ANT-1 and an expression vector for cyclophilin D, a component of the PT pore on the matrix side which directly interacts with ANT-1 (Woodfield et al., 1998). Quantification of apoptosis induction revealed a more than 2-fold reduced cell death by ANT-1 when cyclophilin D was cotransfected (figure 7B).

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Claims

- A method for the inhibition of apoptosis, comprising contacting a cell
 with an effective amount of a substance capable of inhibiting the
 activity of adenine nucleotide translocase-1 (ANT-1).
- 2. The method of claim 1, wherein said cell is a mammalian cell.
- 3. The method of claim 2, wherein said cell is associated with a pathogenic disorder.
- 4. The method of claim 1, wherein the activity of ANT-1 is inhibited on the nucleic acid level.
- 5. The method of claim 4, wherein the inhibition is effected by reducing ANT-1 gene expression.
- The method of claim 4, wherein the activity of the endogenous ANT-1 promoter is reduced.
- 7. The method of claim 1, wherein the activity of ANT-1 is inhibited on the protein level.
- 8. The method of claim 7, wherein the inhibition is effected by adding ANT-1 protein antagonists.
- 9. The method of claim 8, wherein the antagonist is cyclophilin D.
- 10. The method of claim 1, wherein an apoptosis-inducing signal transduction pathway is inhibited, said pathway being activated by ANT-1.
- 11. A method for the treatment of diseases associated with excessive

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apoptosis, comprising the step of administering to a subject in need thereof a pharmaceutically effective amount of a substance capable of inhibiting the activity of adenine nucleotide translocase (ANT-1).

- 12. The method of claim 11, wherein the disease is a degenerative disease.
- 13. The method of claim 12, wherein the disease is dilated cardiomyopathy.
- 14. A method for identifying substances suitable for apoptosis inhibition comprising the step of determining the capability of a test substance to inhibit the activity of ANT-1.
- 15. The method of claim 14, wherein the capability of a test substance to bind ANT-1 or a domain thereof is determined.
- 16. The method of claim 14, wherein the capability of a test substance to bind the N-terminal domain of ANT-1 is determined.
- 17. The method of claim 14, wherein the capability of a test substance to inhibit the binding of ANT-1 to natural binding partners thereof is determined.
- 18. The method of claim 14, which is carried out as a high-throughput assay.
- 19. The method of claim 18, comprising a parallel determination of at least96 test compounds.
- 20. The method of claim 14, which is carried out as a cell-based assay.
- 21. The method of claim 20, which is carried out as an assay using ANT-1-containing cell fractions or ANT-1-containing whole cells.

- 22. The method of claim 14, which is carried out as a molecular-based assay using an isolated protein selected from ANT-1 or a domain thereof.
- 23. The method of claim 22, wherein a recombinant protein is used.
- 24. The method of claim 14, wherein the determining step comprises the measurement of apoptosis induction.
- 25. The method of claim 24, wherein the apoptosis induction is measured by a parameter selected from the group consisting of DNA fragmentation, caspase activation or characteristic alterations in cell morphology.
- 26. A pharmaceutical composition comprising as an active agent an inhibitor of ANT-1 activity, optionally together with pharmaceutic ally acceptable diluents, carriers or adjuvants.
- 27. The pharmaceutical composition of claim 26 for use in the treatment of diseases associated with excessive apoptosis.
- 28. The composition of claim 27 for use in the treatment of human diseases.
- 29. The composition of claim 28 for use in the treatment of dilated cardiomyopathy.
- 30. A method for the diagnosis of an apoptotic process in a degenerative disease or a predisposition therefor comprising detecting the ANT-1 expression in a sample from tissue and/or body fluids of a subject to be tested, wherein elevated ANT-1 expression is indicative for an apoptotic process occuring in a degenerative disease or a predisposition therefor.

31. The method of claim 30, wherein the degenerative disease is dilated cardiomyopathy.

Abstract

The present invention refers to the use of adenine nucleotide translocase-1 (ANT-1), a central component of the permeability transition pore in mitochondria, as a drug target, particularly for the treatment of dilated cardiomyopathy. Further, an assay for the detection of pharmacologically active substances is disclosed which inhibit ANT-1 activity. In addition, ANT-1 expression is claimed as a diagnostic marker for dilated cardiomyopathy.

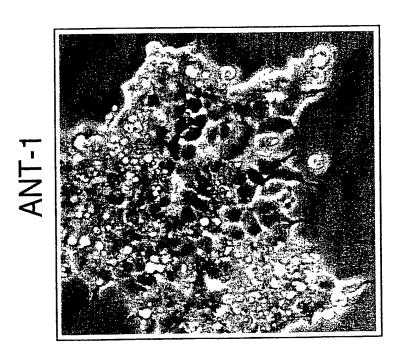
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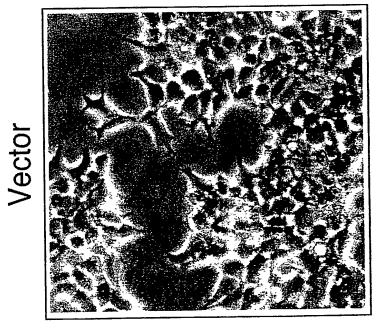
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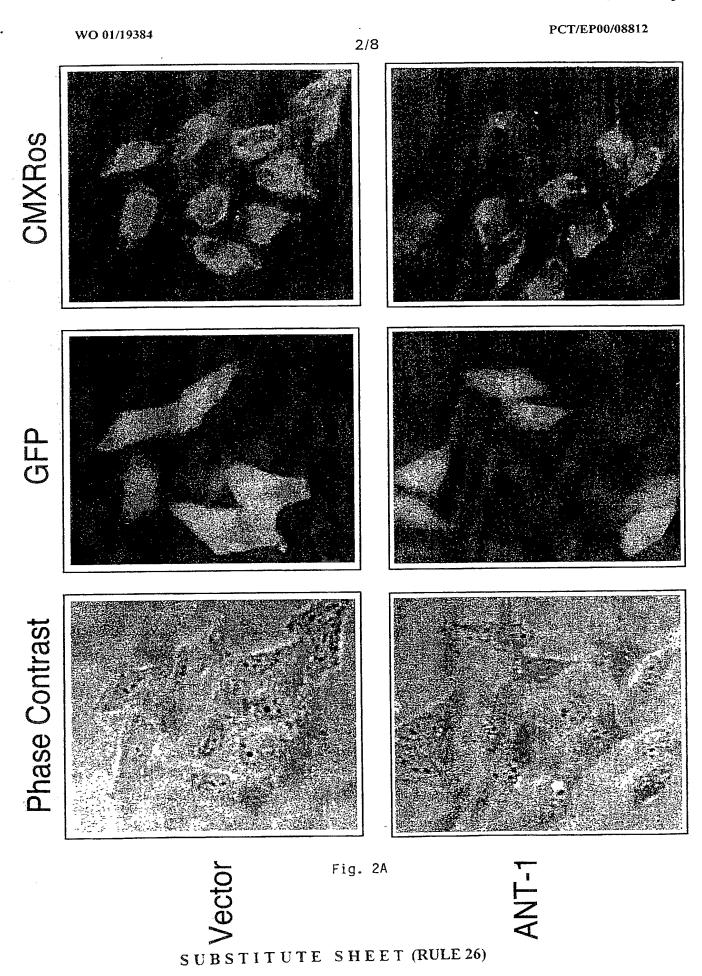
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Fig.1







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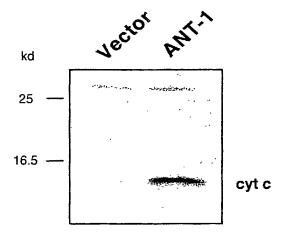


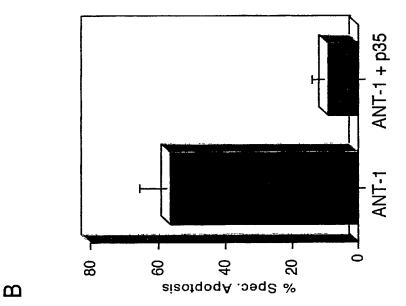
Fig. 2B

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Fig. 3



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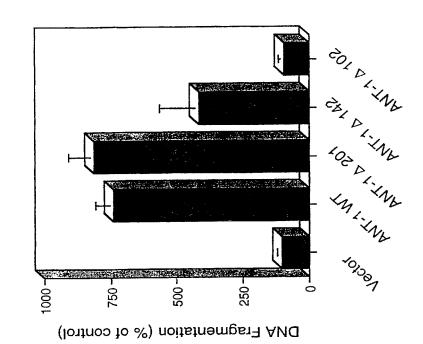
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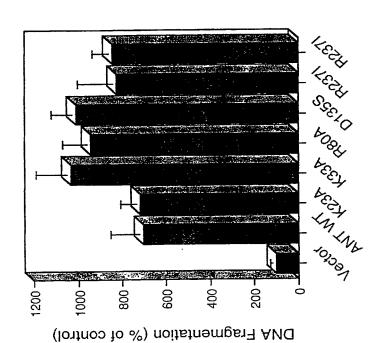
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Fig. 4



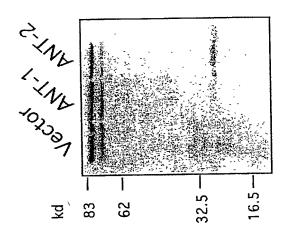


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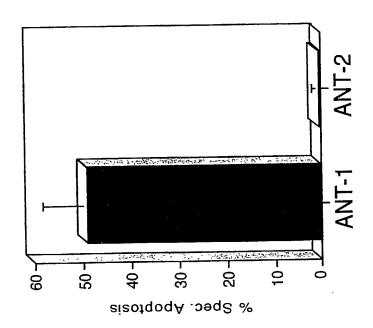
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Fig. 5



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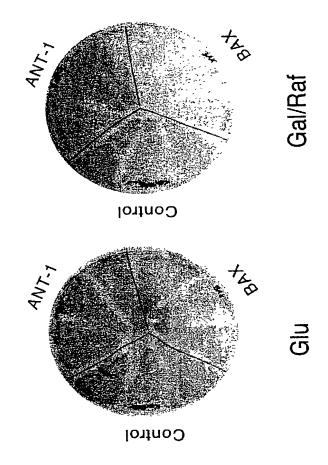


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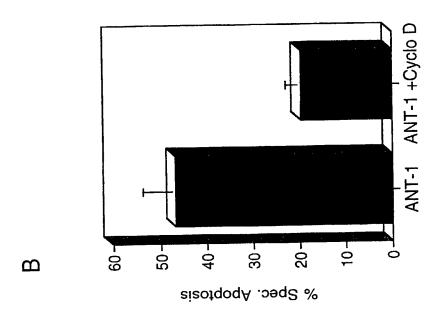
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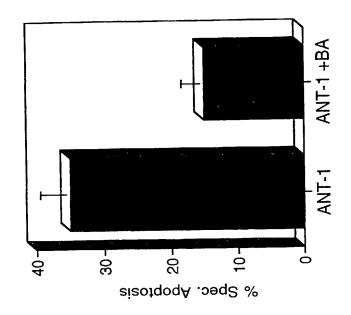
Fig. 6



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Fig. 7





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